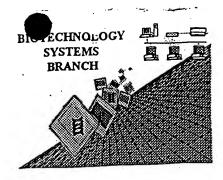
re-run

## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable

Application Serial Number: 09/762, 86/

Source: PG/09

Date Processed by STIC: 2/27/200/

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

Contact

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## **Checker Version 3.0**

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

09/762,867

insert closing angle broadlet um after each response SEQUENCE LISTING <1107Takahashi, Hideaki Sekino, Masachi 1207Method for isolating satellite sequences <130 201487/1060 <140 <141 <150 PCT/JP99/03551 <151 1999-01-07 <150 JP 10-232153 <151 1998-08-18 <160 32 <170 PatentIn Ver. 2.0 <210 1 <211 82 <212 DNA <213 Haliotis discus discus <400 1 cacacacaca cacacacaca cacacacaca cacacacaca cacacacaca cacacacaca 4 < 211 14 < 212 DNA < 213 Haliotis discus discus < 400 4 cacacacaca @ @ @ 14 <210 5 <211 32 <212 DNA <213 Haliotis discus discus <400 5 cacacacaca <210 6 <211 18 <212 DNA <213 Haliotis discus discus <400 6 gagacagaca @ @ 18 <210 7 <211 23 <212 DNA <213 Haliotis discus discus <400 7 caaagaaccc 8 <211 46 <212 DNA <213 Haliotis discus discus <400 8 cgcacgcacg cacgcacgca cgcacgcacg cacgcatgca cccaca @ @ @ @ @ @ @ @ @ 46 <210 9 <211 25 <212 DNA <213 Haliotis discus discus <400 9 cteteteaac aacaacacac 10 <211 50 <212 DNA <213 Haliotis discus discus <400 10 cacacacaca cacacacaca cacacacaca cacacacaca

sample of submitted file major format errors-see coverskeet

Fy I Please review the

Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

> **Does Not Comply** Corrected Diskette Needed